

In the Specification:

Please replace the sequence listing which was filed on October 24, 2002 with the revised sequence listing provided herewith.

Please replace the paragraph found at page 7, lines 10-13, with the following paragraph:

-- *Figure 1* shows the nucleotide sequence (SEQ ID NO:1) and corresponding amino acid sequence (SEQ ID NO:2) of the V_H-region from HuMAb 14.1. CDR regions are also delineated. The germline segments utilized for the HuMAb 14.1 V_H-region include V_H 3-30.3 (SEQ ID NO:9), D 6-13 (SEQ ID NO:12), and JH4b (SEQ ID NO:13). --

Please replace the paragraph found at page 7, lines 15-17, with the following paragraph:

-- *Figure 2* shows the nucleotide sequence (SEQ ID NO:3) and corresponding amino acid sequence (SEQ ID NO:4) of the V_K-region from HuMAb 14.1. CDR regions are also delineated. The germline segments utilized for the HuMAb 14.1 V_K-region include V_K L18 (SEQ ID NO:10) and JK3 (SEQ ID NO:14). --

Please replace the paragraph found at page 7, lines 19-22, with the following paragraph:

-- *Figure 3* shows the nucleotide sequence (SEQ ID NO:5) and corresponding amino acid sequence (SEQ ID NO:6) of the V_H-region from HuMAb 8D2. CDR regions are also delineated. The germline segments utilized for the HuMAb 8D2 V_H-region include V_H 3-30.3 (SEQ ID NO:9), D 7-27 (SEQ ID NO:15), and JH3b (SEQ ID NO:16). --

Please replace the paragraph found at page 7, lines 24-26, with the following paragraph:

-- *Figure 4* shows the nucleotide sequence (SEQ ID NO:7) and corresponding amino acid sequence (SEQ ID NO:8) of the V_K-region from HuMAb 8D2. CDR regions are also delineated. The germline segments utilized for the HuMAb 8D2 V_K-region include V_K A27 (SEQ ID NO:11) and JK2 (SEQ ID NO:17). --